

RAW SEQUENCE LISTING

DATE: 04/22/2002

PATENT APPLICATION: US/09/913,345

TIME: 15:51:17

Input Set : A:\Sequence Listing 5034.ST25.txt Output Set: N:\CRF3\04222002\I913345.raw

```
3 <110> APPLICANT: Garssen, Gerrit J.
             Jacobs, Jorg G.
             Langeveld, Joannes P.M.
             Smits, Marinus A.
                                                                    ENTERED
             van Keulen, Lucien J.M.
             Schreuder, Bram E.C.
             Bossers, Alexander
     11 <120> TITLE OF INVENTION: Prion Test
     13 <130> FILE REFERENCE: 2183-5034US
     15 <140> CURRENT APPLICATION NUMBER: US 09/913,345
C--> 16 <141> CURRENT FILING DATE: 2002-03-25
     18 <150> PRIOR APPLICATION NUMBER: PCT/NL00/00079
     19 <151> PRIOR FILING DATE: 2000-02-09
     21 <150> PRIOR APPLICATION NUMBER: EP 99200391.3
     22 <151> PRIOR FILING DATE: 1999-02-11
     24 <160 > NUMBER OF SEQ ID NOS: 30
     26 <170> SOFTWARE: PatentIn version 3.1
     28 <210> SEQ ID NO: 1
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     43 Thr Gly Gly Ser Arg Tyr Pro Gly Gln Gly Ser Pro Gly Gly Asn Arg
     47 Tyr Pro Pro Gln Gly Gly Gly Gly Trp Gly Gln Pro His Gly Gly
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     55~\mathrm{Trp} Gly Gln Pro His Gly Gly Gly Trp Gly Gln Gly Gly Gly Thr His
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     59 Ser Gln Trp Asn Lys Pro Ser Lys Pro Lys Thr Asn Met Lys His Met
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     63 Ala Gly Ala Ala Ala Gly Ala Val Val Gly Gly Leu Gly Gly Tyr
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     67 Met. Leu Gly Ser Ala Met Ser Arg Pro Ile Ile His Phe Gly Ser Asp
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     71 Tyr Glu Asp Arg Tyr Tyr Arg Glu Asn Met His Arg Tyr Pro Asn Gln
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75 Val Tyr Tyr Arg Pro Met Asp Glu Tyr Ser Asn Gln Asn Asn Phe Val

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165 170 175 79 His Asp Cys Val Asn Ile Thr Ile Lys Gln His Thr Val Thr Thr 185 180 83 Thr Lys Gly Glu Asn Phe Thr Glu Thr Asp Val Lys Met Met Glu Arg 195 200 8' Val Val Glu Gln Met Cys Ile Thr Gln Tyr Glu Arg Glu Ser Gln Ala 220 215 8ส 210 91 Tyr Tyr Gln Arg Gly Ser Ser Met Val Leu Phe Ser Ser Pro Pro Val 92 225 230 95 Ile Leu Leu Ile Ser Phe Leu Ile Phe Leu Ile Val Gly 96 250 245 99 < 210 > SEQ ID NO: 2100 <211> LENGTH: 252 101 <212> TYPE: PRT 102 <213> ORGANISM: Oryctolagus cuniculus 104 <400> SEQUENCE: 2 106 Met Ala His Leu Gly Tyr Trp Met Leu Leu Phe Val Ala Thr Trp 1.0 110 Ser Asp Val Gly Leu Cys Lys Lys Arg Pro Lys Pro Gly Gly Gly Trp 20 25 114 Asn Thr Gly Gly Ser Arg Tyr Pro Gly Gln Ser Ser Pro Gly Gly Asn 35 40 118 Arg Tyr Pro Pro Gln Gly Gly Gly Trp Gly Gln Pro His Gly Gly 55 122 Trp Gly Gln Pro His Gly Gly Gly Trp Gly Gln Pro His Gly Gly 70 75 126 Trp Gly Gln Pro His Gly Gly Gly Trp Gly Gln Gly Gly Thr His Asn 85 90 130 Gln Trp Gly Lys Pro Ser Lys Pro Lys Thr Ser Met Lys His Val Ala 100 105 131 134 Gly Ala Ala Ala Gly Ala Val Val Gly Gly Leu Gly Gly Tyr Met 135 115 120 138 Leu Gly Ser Ala Met Ser Arg Pro Leu Ile His Phe Gly Asn Asp Tyr 130 135 142 Glu Asp Arg Tyr Tyr Arg Glu Asn Met Tyr Arg Tyr Pro Asn Gln Val 155 150 146 Tyr Tyr Arg Pro Val Asp Gln Tyr Ser Asn Gln Asn Ser Phe Val His 165 170 175 150 Asp Cys Val Asn Ile Thr Val Lys Gln His Thr Val Thr Thr Thr 185 190 180 154 Lys Gly Glu Asn Phe Thr Glu Thr Asp Ile Lys Ile Met Glu Arg Val 155 195 200 158 Val Glu Gln Met Cys Ile Thr Gln Tyr Gln Glu Ser Gln Ala Ala 220 210 215 162 Tyr Gln Arg Ala Ala Gly Val Leu Leu Phe Ser Ser Pro Pro Val Ile 230 166 Leu Leu Ile Ser Phe Leu Ile Phe Leu Ile Val Gly 245 170 < 210 > SEQ ID NO: 3

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171 <211> LENGTH: 254 172 <212> TYPE: PRT 173 <213 > ORGANISM: Syrian golden hamster 175 <400 > SEQUENCE: 3 177 Met Ala Asn Leu Ser Tyr Trp Leu Leu Ala Leu Phe Val Ala Met Trp 1.0 181 Thr Asp Val Gly Leu Cys Lys Lys Arg Pro Lys Pro Gly Gly Trp Asn 20 25 185 Thr Gly Gly Ser Arg Tyr Pro Gly Gln Gly Ser Pro Gly Gly Asn Arg 186 35 40 189 Tyr Pro Pro Gln Gly Gly Gly Thr Trp Gly Gln Pro His Gly Gly Gly 55 193 Trp Gly Gln Pro His Gly Gly Gly Trp Gly Gln Pro His Gly Gly 70 75 197 Trp Gly Gln Pro His Gly Gly Gly Trp Gly Gln Gly Gly Gly Thr His 85 90 201 Asn Gln Trp Asn Lys Pro Ser Lys Pro Lys Thr Asn Met Lys His Met 202 100 105 205 Ala Gly Ala Ala Ala Gly Ala Val Val Gly Gly Leu Gly Gly Tyr 115 120 209 Met Leu Gly Ser Ala Met Ser Arg Pro Met Met His Phe Gly Asn Asp 135 213 Trp Glu Asp Arg Tyr Tyr Arg Glu Asn Met Asn Arg Tyr Pro Asn Gln 155 150 217 Val Tyr Tyr Arg Pro Val Asp Gln Tyr Asn Asn Gln Asn Asn Phe Val 165 170 221 His Asp Cys Val Asn Ile Thr Ile Lys Gln His Thr Val Thr Thr 180 185 225 Thr Lys Gly Glu Asn Phe Thr Glu Thr Asp Ile Lys Ile Met Glu Arg 195 200 229 Val Val Glu Gln Met Cys Thr Thr Gln Tyr Gln Lys Glu Ser Gln Ala 215 233 Tyr Tyr Asp Gly Arg Arg Ser Ser Ala Val Leu Phe Ser Ser Pro Pro 234 225 235 230 237 Val Ile Leu Leu Ile Ser Phe Leu Ile Phe Leu Met Val Gly 238 245 241 <210> SEQ ID NO: 4 242 <211> LENGTH: 254 243 <212> TYPE: PRT 244 <213> ORGANISM: Murinae gen. sp. 246 <400> SEQUENCE: 4 248 Met Ala Asn Leu Gly Tyr Trp Leu Leu Ala Leu Phe Val Thr Met Trp 252 Thr Asp Val Gly Leu Cys Lys Lys Arg Pro Lys Pro Gly Gly Trp Asn 20 256 Thr Gly Gly Ser Arg Tyr Pro Gly Gln Gly Ser Pro Gly Gly Asn Arg 35 4.0 260 Tyr Pro Pro Gln Gly Gly Thr Trp Gly Gln Pro His Gly Gly Gly Trp 261 50 55

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264 Glv Gln Pro His Gly Gly Ser Trp Gly Gln Pro His Gly Gly Ser Trp 265 65 268 Gly Gln Pro His Gly Gly Gly Trp Gly Gln Gly Gly Gly Thr His Asn 85 90 272 Gln Trp Asn Lys Pro Ser Lys Pro Lys Thr Asn Leu Lys His Val Ala 273 100 105 110 276 Gly Ala Ala Ala Gly Ala Val Val Gly Gly Leu Gly Gly Tyr Met 120 125 180 Leu Gly Ser Ala Met Ser Arg Pro Met Ile His Phe Gly Asn Asp Trp 135 140 281 130  $\pm 84~{
m Glu}$  Asp Arg Tyr Tyr Arg Glu Asn Met Tyr Arg Tyr Pro Asn Gln Val 155 150 288 Tyr Tyr Arg Pro Val Asp Gln Tyr Ser Asn Gln Asn Asn Phe Val His 170 165 292 Asp Cys Val Asn Ile Thr Ile Lys Gln His Thr Val Ihr Ihr Thr Thr 180 185 296 Lys Gly Glu Asn Phe Thr Glu Thr Asp Val Lys Met Met Glu Arg Val 195 200 300 Val Glu Gln Met Cys Val Thr Gln Tyr Gln Lys Glu Ser Gln Ala Tyr 210 215 220 304 Tyr Asp Gly Arg Arg Ser Ser Ser Thr Val Leu Phe Ser Ser Pro Pro 235 305 225 230 308 Val Ile Leu Leu Ile Ser Phe Leu Ile Phe Leu Ile Val Gly 245 312 < 210 > SEQ ID NO: 5313 <211> LENGTH: 256 314 <212> TYPE: PRT 315 <213> ORGANISM: Bos taurus 317 <400> SEQUENCE: 5 319 Met Val Lys Ser His Ile Gly Ser Trp Ile Leu Val Leu Phe Val Ala 5 10 323 Met Trp Ser Asp Val Gly Leu Cys Lys Lys Arg Pro Lys Pro Gly Gly 327 Gly Trp Asn Thr Gly Gly Ser Arg Tyr Pro Gly Gln Gly Ser Pro Gly 3.5 40 331 Gly Asn Arg Tyr Pro Pro Gln Gly Gly Gly Gly Trp Gly Gln Pro His 335 Gly Gly Gly Trp Gly Gln Pro His Gly Gly Gly Trp Gly Gln Pro His 7.0 75 339 Gly Gly Gly Trp Gly Gln Pro His Gly Gly Gly Gly Trp Gly Gln Gly 90 8.5 343 Gly Thr His Gly Gln Trp Asn Lys Pro Ser Lys Pro Lys Thr Asn Met 100 105 347 Lys His Val Ala Gly Ala Ala Ala Gly Ala Val Gly Gly Leu 115 120 351 Gly Gly Tyr Met Leu Gly Ser Ala Met Ser Arg Pro Leu Ile His Phe 135 140 352 130 355 Gly Ser Asp Tyr Glu Asp Arg Tyr Tyr Arg Glu Asn Met His Arg Tyr 155

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Input Set : A:\Sequence Listing 5034.ST25.txt
Output Set : N:\CRF3\04222002\1913345.raw

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350 353	Λen	Dha	Val	Hic	165	Cvs	Val	Δen	Ile	170 Thr	Val	Ivs	Glu	His	175 Thr	Val
364	ASII	rne	vai	180	KSP	CYS	Vai	NSII	185	1111	vai	LIS	JIU	190	1111	VQI
	Thr	Thr		Thr	Lys	Gly	Glu		Phe	Thr	Glu	Thr		Ile	Lys	Met
368 371	Met	Glu	195 Arg	Val	Val	Glu	Gln	200 Met	Cys	Lle	Thr	Gln	205 Tur	Gln	Ara	Glu
372	1100	210	*** 9	7 4 2 2	, 41	314	215		$\cup_I D$			220	- 1 -	92	,	
		Gln	Ala	Tyr	Tyr	Gln 230	Arg	Gly	Ala	Ser	Val 235	Ile	Leu	Phe	Ser	Ser 240
	225 Prio	Pro	Val	Ile	Leu		Ile	Ser	Phe	Leu		Phe	Leu	Ile	Val	
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		Trp	Ser	Asp	_	Glv	Leu	Cys	Lys		Arq	Pro	Lys	Pro	_	Gly
395		1		20		1		- 1 -	25	1 -	,		4	30	1	1
398	Gly	Prp	Asn	Thr	Gly	Gly	Ser	Arg	Tyr	Pro	Gly	Gln	Gly	Ser	Pro	Gly
399			35					40	_		_		45			•
402	Gly	Asn 50	Arg	Tyr	Pro	Pro	Gln 55	Gly	Gly	Gly	Gly	Trp 60	GLy	GIn	Pro	His
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411	.a.,	_			85	_	_		_	90	_	_	_	<b>~</b> 1	95	
414	GLY	Ser	HIS	Ser 100	GIn	Trp	Asn	Lys	Pro 105	ser	Lys	Pro	Lys	110	Asn	мет
	Lys	His	Val		Gly	Ala	Ala	Ala		Gly	Ala	Val	Val		Gly	Leu
419			115		_	_ 7		120		~	_	_	125	- 1		<b>5</b> 1
422	Gly	G1y 130	Tyr	Met	Leu	GIY	Ser 135	Ala	Met	ser	Arg	Pro 140	Leu	He	HIS	Pne
	_	Asn	Asp	Tyr	Glu		Arg	Tyr	Tyr	Arg		Asn	Met	Tyr	Arg	
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431	PIO	ASII	GIII	Val	165	1 7 1	Arg	FIO	Vai	170	лгу	тут	561	ASII	175	ASII
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435				180		_	_		185		_			190	_	1
438 439	Thr	Thr	Thr 195	Thr	Lys	Gly	Glu	Asn 200	Phe	Thr	Glu	Thr	Asp 205	Ile	Lys	He
442	Met		Arg	Val	Val	Glu		Met	Cys	Ile	Thr		Tyr	Gln	Arg	Glu
443		210	_				215			_		220		_,	.=	.~
		Gln	Ala	Tyr	Tyr		Arg	Gly	Ala	Ser		Ile	Leu	Phe	Ser	
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VERIFICATION SUMMARY

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PATENT APPLICATION: US/09/913,345 TIME: 15:51:18

Input Set : A:\Sequence Listing 5034.ST25.txt

Output Set: N:\CRF3\04222002\I913345.raw

L:16 M:271 C: Current Filing Date differs, Replaced Current Filing Date